AMENDMENTS TO THE SPECIFICATION

Please amend the specification as follows:

Please amend the paragraph on page 12, lines 24-30:

For example, synthesis of the first cDNA strand may be carried out in the presence of primers having appropriate restriction sites in order to allow a subsequent cloning in the proper orientation with respect to the expression vector promoter. As restriction sites, any known restriction site may be used. As a primer, for instance the following primer, 50 nucleotides long may be used: (SEQ ID NO: 181)

Please amend the paragraph bridging pages 12 and 13:

After two-strand synthesis, the cohesive ends of the double stranded cDNA are filled (blunt end) and the cDNA ends are then ligated using a suitable DNA adaptor sequence. The DNA adaptor sequence should contain a restriction site which should be different from the restriction site used in the primer for the synthesis of the first cDNA strand. The DNA adaptor may comprise for example complementary 9- or 13-mer oligonucleotides, whose ends represent the cohesive end of a restriction site. These ends may be for example a EcoRI-site:

B3 Cond 5' XXXXXGGCACGAG 3' (SEQ ID NO: 182)

3' XCCGTGCTC 5'

Please amend the paragraphs on page 24, lines 3-32:

pKS+\(\Delta\)NotI is cleaved with PstI and BamHI and the DNA oligonucleotide, synthesized from the pK3/pK4 primer pair described below, is ligated in the opened plasmid. The pKS+neu plasmid thus prepared contains between PstI and BamHI restriction sites, the following novel restriction sites NotI, StuI, SfiI and NcoI (i.e. PstI-NotI-StuI-SfiI-NcoI-BamHI)

- 5'-GCGGCCGCAAGGCCTCCATGGCCG-3' PK3 (SEQ ID NO: 183)
- 5'-GATCCGGCCATGGAGGCCTTGCGGCCGCTGCA-3' PK4 (SEQ ID NO: 184)

The URA3 gene of S.cerevisiae is amplified via PCR, by use of the primer-pair PK9 and PK10, described below, and an Ycplac33 vector DNA (Gietz, R. D. and Sugino, A. (1988) Gene 74: 527-534) as matrix. The amplified DNA is cleaved with BamHI and NotI and subsequently inserted in pKS+neu which has been cleaved by BamHI and NotI. The plasmid thus obtained is named pPK9/10.

..NotI..
5'-ATCTGCAGCGGCCG<u>CAAACATGAGAATTGGGTAATAACTG</u>-3' PK9

PstI

(SEQ ID

NO:185)

..SfiI..

5'-ATGGATCCGGCCATGGAGGCCTTCAAGAATTAGCTTTTCAATTCATC-3'

BamH1

PK10 (SEQ ID

NO: 186)

187)

2) Preparation of the deletion cassette

The 5'-region of ORF YML114c was amplified by PCR using genomic DNA of S.cerevisiae as template and both primers YML114c-Asp718 and YLM114c-EcoRI, described below.

YML114c-Asp718: 5'-GCTGGTACCCGTCGGTCTCTTTACC-3'(SEQ ID NO:

YLM114c-EcoRI: 5'-TTGGAATTCATTGCCCTTTATGAGTCC-3'(SEQ ID NO: 188)

Please amend the paragraph on page 25, lines 1-9:

The 3'region of ORF YML114c was amplified by PCR using genomic DNA of S.cerevisiae as template and both primers YML114c-BamHI and YLM114c-SacI, described below.

YML114c-BamHI:5'-ATCGGATCCGCCAACAATGACAGCG-3'(SEQ ID NO: 189)

YLM114c-SacI: 5'-GTTGAGCTCTGAGCGTTTGTCCTTG-3'(SEQ ID NO: 190)

The PCR fragment was subsequently cut with BamHI and SacI. The resulting 535bp fragment was inserted in plasmid pYML114c-A linearized with BamHI and SacI generating pYML114c-B.

TABLE 2: Primers used for gene deletions

Gene deletion	ns on chromosome 13
Name	Sequence 5'-3'
YDR472w-S1	ATG TCT CAA AGA ATA ATT CAA CCA AGC GCA TCT GAC
	CAA CCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 167)
YDR472w-S2	AGC CAA ATC TCA AAC CTT CCC TGT CAA GCA CTT GCC
	TGT CGC ATA GGC CAC TAG TGG ATG TG (SEQ ID NO:
	177)
YDR499w-S1	ATG AGA CGA GAA ACG GTG GGT GAA TTT TCT TCA GAT
	GAC GCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 95)
YDR499w-S2	CGT ACT TTA CTT GCA TTA TTC TCC CCG TTC TTT TAT
	TCA AGC ATA GGC CAC TAG TGG ATG TG (SEQ ID NO:
IDD 040 = .01	175)
YMR049c-S1	CAG ACT ATT GAT TAC TTT ATG ACC GGT TAG TTT CTT
mm040 G0	TAG TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 170)
YMR049c-S2	TCT GTT CTA ACA TAA CTA GGT CAA TGA TGG CTA AGA
	ACA AGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO: 128)
YMR134w-S1	GCA AAG TGT GGT ATA GAA AAA GAA CCA AAG GCC GGT
III(IJW DI	ATG TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 139)
YMR134w-S2	TGT GTG TGT GCC TAC CTG CAT GTA TGC ATT TAG CAA
114(134" 52	TTG AGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
	150)
YML023c-S1	CAC GCA ATG GTG CAC ATT ATT TTG TTG AAC TCA CTG
	AGA ACA GCT GAA GCT TCG TAC GC (SEQ ID NO: 159)
YML023c-S2	ATT AGT TAC TTA TTC TAT AAT TAC ACT TTT ATC ATG
	AAC GGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
	173)
YML049c-S1	AAT TCC TGC TCA TTC AAG GAA AGT CTC AGG AAA TTT
	TCA CCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 106)
YML049c-S2	ACT CCT GCA TCG GAC ACT TCG TCG ATC TGG AAG CAG
	GGT CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO: 19)
YML077w-S1	ATG GGG ATA TAT TCA TTT TGG ATC TTT GAT AGG CAT
	TGT ACA GCT GAA GCT TCG TAC GC (SEQ ID NO: 28)
YML077w-S2	TTC TAT TGG TGA TCT TTC TTG TCC CTT GAC CTC TCA
	TTT CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO: 39)
YML093w-S1	GCT AAC TTA AAT ATG GCA AAA AAG AAA TCT AAG AGC
	AGA TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 50)
YML093w-S2	CAA AGG ATC AAT AAC TTG GCC TGG CTT AGT CAT GAT
	TCT CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO: 61)
YML114c-S1	AAC GTG TAA TTG AGG GAC TCA TAA AGG GCA ATG ACT
	TCC ACA GCT GAA GCT TCG TAC GC (SEQ ID NO: 71)

YML114c-S2	GAC	TTG	TAG	TAG	CAT	CGA	TAT	TGG	TTG	TGT	TAT	GTG
	CTA	C GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u> §	SEQ ID	NO:	<u>81)</u>
YML127w-S1	CCG	CTA	AAT	GGT	ACT	CCA	GTA	AGC	GAG	GCA	CCC	GCC
	ACA	A CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ II	NO:	84)	
YML127w-S2	ATA	ACC	CCG	ACG	TGT	TTT	CCA	TGT	ATT	CAG	ACA	ATG
	CTA	A GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u> §	SEQ ID	NO:	<u>85)</u>



Gene deletions	on (chro	nosoi	me 13	3							
Name	Sequ	ence	5'-	-3'								
YMR032w-S1										CCC		CGA
	+							-		D NO:		
YMR032w-S2		AAA								CGA		
	+									SEQ II		
YMR093w-S1	ATG		ACT							ACT		AAG
								=		D NO:		ma.
YMR093w-S2		CAC			AGT					GTA		
mm121 = 01	CTT	CGC		CCT						SEQ II		
YMR131c-S1										ACA D NO:		WII
YMR131c-S2	GGT									ATA		ጥርር
IMRISIC-52										SEQ II		
YMR185w-S1	ATC									AGA		
1120311 52							TAC			D NO:		
YMR185w-S2	GTA	ATG								GTT		AGC
	TTG	T GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (SEQ II	ONO:	93)
YMR212c-S1	CCT	CTT	GAA	CTT	AAA	GAA	TGT	AAA	TCT	TCA	ттт	GCG
	TCT	T CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ I	D NO:	94)	
YMR212c-S2	CGG	ATG	ATG	TTC	ACA	CCA	AAA	CAT	CAG	AAA	CTG	GTC
	AAT	C GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (SEQ II	ONO:	96)
YMR213w-S1	ATA	CGT	GAA	AGG	CGG	TGT	ATG	GAC	CAA	TGT	GGA	GGA
	TCA	G CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ I	D NO:	<u>97)</u>	
YMR213w-S2	GCT	GTA	ACT	GTT	CAA	TAG	ACT	CCA	CTT	TTG	ATT	GGA
	TCG	A GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (SEQ II	NO:	98)
YMR218c-S1	GAC	TCA	AAT						ACT	CTA	CAA	CTT
	 	CCA					TAC			D NO:		
YMR218c-S2		GGC		_						GGT		
	GCT	T GC	A'1' <i>P</i>	A GG(C CA	C TA	AG T'	∃G A	TC T	'G <u>(s</u>	EQ ID	NO:
YMR281w-S1	100)	AAG	222	ልርጥ	Тλλ	ATC	AAG	ATG	ጥሞር	AGG	ССТ	ACA
114120111 51	1									D NO:		
YMR281w-S2	 		i					_		AGT		AGC
	TTG									G (s		
	102)											
YMR288w-S1	GAA	AAC	CTG	CAG	AAA	GAA	GCT	GCA	CGT	ATT	GGT	GAG
	AAC	G CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ I	D NO:	103)	
YMR288w-S2	CCA	AAC	CTT	CTA	AAA	TAC	GCA	TAA	TAG	CAT	GTG	GTG
		TGC	ATA	4 GG	C CA	C TA	AG TO	GG A	TC I	'G <u>(s</u>	EQ ID	NO:
17m200 - 61	104)				~~~		055	Mwz	000	000	mmm	003
YMR290c-S1										GTT		CCA
VMP 200 g C2	+									D NO:		CCT
YMR290c-S2										AGT		
	107)	CGC	ATF	1 GG(L CA	.C TF	1G T	A DE	1 C 1	'G <u>(s</u>	EQ ID	NO:
L	1 1 0 /)											

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YMR211w-S1	AGA	GAG	CAA	ACC	ATT	TGA	CTA	CTO	CAAT	TCT	TCA	ATA
	TAC	A CA	GCT	GAA	GCT	TCG	TAC	GC .	(SEQ	ID NO:	108)	_
YMR211w-S2	ATT	TCA	ATC	ATC	TTA	CTC	CGT	GAZ	A TC	GGT	TCG	GAA
	TGA	TGC	ATA	GG	C CA	C TA	AG T	GG A	ATC '	rg <u>(s</u>	EQ ID	NO:
	109)											

13/2 Usrd

Gene deletions	s on chromosome 4
Name	Sequence 5'-3'
YDR196c-S1	ATG CTT ATG ATC AAA TTG TGT TAT ACT TCA AGG ACA
	AAA TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 110)
YDR196c-S2	TTT CAA TCT GTT CGT ATA AGT CAA CCA ATG TGC TGT
	TAT TGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
	111)
YDR299w-S1	ATG GAA AAA TCA CTA GCG GAT CAA ATT TCC GAT ATC
	GCC ACA GCT GAA GCT TCG TAC GC (SEQ ID NO: 112)
YDR299w-S2	CAA AGA TTT GGA TAT CAT CGT TTT TAA CAG CCT CTA
	ATT CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
	113)
YDR365c-S1	CTG GAG AGA ACC CAA AGA AGG AAG GTG TAG ATG CTA
	GGT TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 114)
YDR365c-S2	TTA GTA TGC TTT TTA TTA ACA GAT TTC AAC TTG CTT
	TTC TGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
	115)
YDR396w-S1	CAG ATA CAC TAT TGT GGT GTA ATC TGG ACC TTG ACT
	GTC TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 116)
YDR396w-S2	TAG AGA AAA CAC TGA ATG ATC TTA GCG ACC GTA CAA
	AAG AGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
	118)
YDR407c-S1	TTC TTA AGC ATT TCC CAA GCT ATG TTG GCC CAT CTA
	AGA TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 119)
YDR407c-S2	AAT AAC AGA CAA GAT AAC GTT TTC AGA GTC GAA CTG
	GAT TGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
	120)
YDR416w-S1	ACT TAC ATG GAA AAG ATA TAT CGA GTA TTG GAA AGA
	GGA GCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 121)
YDR416w-S2	TCA AAT ATC TAG TTC TAT TTC ATC TGG ATT AAT CGA
	ATA TGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
	122)
YDR449c-S1	CAC ATC ACC GAT TTC TAA TAA TGT CGA AGA CAA GAT
	ACT ACA GCT GAA GCT TCG TAC GC (SEQ ID NO: 123)
YDR449c-S2	ATA ATT AAA TCT AGA ATT TTA TAC CTA GGA TCA TCT
	TCT GGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
VDD141 a - C1	TTC GTA ATC TTT GAA TTC TGC GAT TTC ATC TAC CAG
YDR141c-S1	
11DD 1 4 1 60	CGC GCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 125)
YDR141c-S2	CAC TAA AGC CCC TTA CAA TTG ACT CAA ATA ATA AAC
	AAC TGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
YDR324c-S1	AAG AAG CCT GAA AAT ACG AAA CAA ACC GGT GAA GAT
IDK324C-SI	
VDD 2045 60	GAC CCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 127)
YDR324c-S2	AAA CACTAA CTT TGG TTG AAT AAA CGC CTT TTG TTT
	GGA GGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
	129)

13 b

YDR325w-S1	GAC	ATT	AAT	ACG	AAA	ATC	TTT	AAC	TCA	GTT	GCT	GAA
	GTA	T CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ II	NO:	130)	
YDR325w-S2	ACC	TCG	CTG	AAA	GAC	TCT	GAA	TCC	TTA	TCT	TCT	TCA
	TCT	A GC	ATA	GG(C CA	C TA	G TO	GG A'	TC TO	G <u>(s</u> i	EQ ID	NO:
	<u>131)</u>											
YDR398w-S1	ATG	GAT	TCT	CCT	GTT	CTA	CAG	TCC	GCT	TAT	GAC	CCA
	TCA	G CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	NO:	132)	
YDR398w-S2	AAC	GTC	ACT	ATA	TCC	GGC	TTC	CTC	CTC	GCC	GTC	GCT
	CTG	C GC	ATP	GG(C CA	C TA	G TO	GG A'	TC TO	G <u>(SI</u>	EQ ID	NO:
	133)											

BL

Gene deletion	s on chromosome 4
Name	Sequence 5'-3'
YDR246w-S1	ATG GCC ATC GAA ACA ATA CTT GTA ATA AAC AAA TC
	GGC GCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 134)
YDR246w-S2	AAC AGG TTA GAT CTT ATA GGC ATT TCC ATT GAG TA
	GAT GGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	135)
YDR236c-S1	CTA AAA TAT TGA ACT TGA CCC TGG CCC CAT AAA AA
	CAT TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 136)
YDR236c-S2	TTG AAG TGT TGA TGT TTACGT GGA CTA TTT ATG TT
	CGT TGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	137)
YDR361c-S2	TTA CCA AGT GGA AAT TTC TGT TTC CAA TTC ATC GA
	ACT TGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
YDR361c-S1	GGT TCA AGC TAT CAA ATT AAA TGA TTT AAA AAA TA
IDRJ01C-BI	GAA GCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 140)
YDR367w-S1	ATC TGC GTA CTT TAT ACA ATC GAT ACC ATT TCC AC
IDRSO/W DI	TGT TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 141)
YDR367w-S2	GTT TTG TTC TAC GTC ATC CCT ATC AAC TAA ATA TT
	GGG GGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	142)
YDR339c-S1	TAT GGG TAA AGC TAA GAA AAC AAG AAA GTT TGG CC
	CGT ACA GCT GAA GCT TCG TAC GC (SEQ ID NO: 143)
YDR339c-S2	TAA AAG ACA TCT GGC AAT TTT TCA ATG ACG TAT GC
	TGA CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	144)
YDR413c-S1	TTC TTT GGT TTA TTC TTC GTT CAT TTT TGG TCA AA
	ATC TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 145)
YDR413c-S2	ACA AAA GAA AGC ACA AGA GTT TAT TAA GGA GCA GG
	AAG GGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	146)
YDR429c-S1	TCT AGA TCT ATC ATT ACA TAC AAG ATT GAA GAC GG
TEND 400	GTC ACA GCT GAA GCT TCG TAC GC (SEQ ID NO: 147)
YDR429c-S2	TTT CTT TGT TTC TAA CGA CAG AAA CTC TTG GAA TG
	GTG CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
YDR468c-S1	GTC ACA ATA CTG CTG GTG ATG ACG ATC AAG AGG AG
IDRIGOCO DI	AAA TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 149)
YDR468c-S2	CAA GAC GAC AAT AAG AAG TCC TAT ACA ACA ATC GT
	GTA TGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	151)
YDR489w-S1	ACT ACC CAC AGA GAT GCA AAT ACA ATA GTG GGT TC
	TCC TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 152)
YDR489w-S2	AGT CGG GCT CAT CTA TCA TGT TTA CGC TAC CTT CT
	TAT CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	178)

cond

YDR527w-S1	ATG	GAC	TTA	CTG	GGC	GAT	ATA	GTG	GAG	AAA	GAT	ACA
	TCT	G CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	NO:	<u>179)</u>	
YDR527w-S2	CCC	CAC	CGC	CTT	GTT	TCC	ATA	ACC	AAA	GTG	CAT	CAA
	TAG	C GC	ATA	GG(C CA	C TA	G TO	G A	rc To	G <u>(S</u> I	EQ ID	NO:
	<u>153)</u>			_								
YDR288w-S1	ATG	AGT	TCT	ATA	GAT	AAT	GAC	AGC	GAT	GTG	GAT	TTA
	ACA	G CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	NO:	154)	
YDR288w-S2	GCC	CAT	GAT	TTC	TTG	CAC	CAA	TTT	TTC	AAG	AGA	CTC
	TAG	T GC	ATA	GG(C CA	С ТА	G TO	G A	rc ro	G (SI	EQ ID	NO:
	155)											

Gene deletions	on	chro	nosor	me 4								
Name	Sequ	ence	5'-	-3 '								
YDR201w-S1	CCC	ATG	тст	GGA	CTA	TTC	AGA	GCA	TCA	TCG	TCA	TCC
	ATA	C CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ II	NO:	156)	
YDR201w-S2	AAA	AGG	GTT	TTC	CGT	TTA	GTT	CCC	GAA	TAT	GAT	GTT
	GAA	A GC	ATA	GG(C CA	C TP	AG TO	GG A'	TC T	G <u>(s</u> :	EQ ID	<u>NO:</u>
	157)											
YDR434w-S1	ATG	TCC	AAT	GCA	AAT	CTA	AGA	AAA	TGG	GTT	GGT	TTT
	TGC	T CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ II	NO:	158)	
YDR434w-S2	TAA	AGG	TAA	ATA	CAC	AGC	TAT	CAT	GTG	CTC	TTG	TGG
	GAA	G GC	ATA	GG(C CA	C TA	G TO	GG A	rc ro	G <u>(s</u> i	O ID	NO:
	160)											
YDR181c-S1	AGG	ATA	AAC	CCA	AAT	GCT	GGA	CAT	CTA	AGG	AAA	TCT
	AAG	T CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	NO:	161)	
YDR181c-S2	TAG	TTG	GGT	TTG	AAT	CGT	TAT	CAC	GGG	AGA	ACA	TTG
	CTT	TGC	ATA	GG(C CA	C TA	AG TO	GG A'	TC T	G <u>(s</u> :	EQ ID	NO:
	162)											
YDR531w-S1	ATG	CCG	CGA	ATT	ACT	CAA	GAG	ATA	TCT	TAC	AAT	TGC
	GAT	T CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	NO:	163)	
YDR531w-S2	AAA	TAA	GCT	ATT	TGC	CCA	ATA	TTG	TTG	GAG	ATG	GCG
	AAT	A GC	ATA	GG(C CA	C TA	AG T	GG A	TC T	G <u>(s</u> :	EQ ID	NO:
	164)											

B6 Const

Gene deletions	on o	chro	noson	me 1	2							
Name	Sequ	ence	5 '-	-3'								
YLR186w-S1	CTA	GTC	ACC	AAG	AAG	AAA	ACC	CGT	AAA	ATC	GTA	GGT
	CAT	G CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	O NO:	165)	
YLR186w-S2	ATA	CAA	AGA	GGA	TGC	CAA	GTA	GAC	TTA	AAC	ACT	ATA
	AAA	TGC	ATA	A GG	C CA	C TP	G T	GG A'	TC T	G (s	EQ ID	NO:
	166)											
YLR215c-S1											AAC	TTT
								GC (
YLR215c-S2	1										GCT	
	1	A GC	ATF	A GG	C CA	.C TA	G TO	GG A'	TC T	G <u>(s</u> :	EQ ID	<u>NO:</u>
	169)											
YLR222c-S1											AAT	ATG
	 							GC <u>(</u>				
YLR222c-S2											ACC	
	1	A GC	ATA	A GG	C CA	C TA	G T	GG A'	rc T	G <u>(s</u>	EQ ID	NO:
WT DO 42 01	174)	mam	000				3.00	CM3	mm »	001	CCM	003
YLR243w-S1	_	_		-							CCT	GCA
	+							GC <u>(</u>				
YLR243w-S2				-	_						TTA	
	1	AGC	A'T'A	A GG	C CA	.C TP	4G 'I'(G A'	rc r	G <u>(S</u>	EQ ID	NO:
YLR272c-S1	171)	000	mcm.	000	3 CM	mmc	mc a	CITIC	mmc	CAA	CTA	3 MM
1LR2/2C-51	1							GC (AII
YLR272c-S2	 											CCM
1LR2/2C-52	1										CCG EQ ID	
	172)	GGC	Alf	1 GG	CCA	.C 12	70 10	JG A	10 1	G <u>(S</u>	EQ ID	NO:
YLR275w-S1	_	TTT	TAT	CAT	GTC	GTA	TGT	TTG	ATC	TTA	ACC	ATT
	ттт	A CA						GC (
YLR275w-S2	 										GTT	TAC
	TTA	T GC	АТА	GGC	CAC	TAG	TGG	ATC	TG (S	SEO II	NO:	4)
YLR276c-S1	CTT		CGG								CTT	_
	GAC	T CA	GCT	GAA	GCT	TCG	TAC	GC (SEO II	NO:	5)	
YLR276c-S2	+										TAT	ATT
	1										NO:	
YLR317w-S1	 					-			785.20		TTT	
	GAT	G CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ II	NO:	7)	
YLR317w-S2	+										TCG	AAA
	1										EQ ID	
	176)					·						
YLR359w-S1	GGC	TAT	TGC	TGA	GAA	GGA	ATT	GGG	CTT	AAC	TGT	TGT
	TAC	A CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	NO:	8)	
YLR359w-S2	+										TGG	AAA
	1										NO:	
YLR373c-S1	•										TGG	
	TGC	C CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ II	NO:	10)	
L												

Bb

YLR373c-S2	CAA	ACA	GAC	TTT	GTT	CCT	TTG	TAT	GTC	CTA	TGG	AAG
	ATA	C GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (S	EQ ID	NO:	11)
YLR424w-S1	GAC	ATG	ACA	TAC	ACT	AAT	GAT	GCC	TTG	AAA	ACT	AGT
	AGC	G CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	NO:	12)	
YLR424w-S2	ATA	GGT	ACT	TTC	TAG	AGG	TCA	AGG	GCC	CAT	AAA	TAA
	ATT	G GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u> 9	EQ ID	NO:	13)

Gene deletion	s on	chro	nosoi	m 1:	2							
Name	Sequ	ence	5 '-	-3'								
YLR437c-S1	ATT	GTG	CAA	GTC	TGT	TAA	AGT	CTT	СТС	TTG	GAT	CCA
	TTA	A CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ I	O NO:	14)	
YLR437c-S2	CAT	CAC	ACA	CTA	ATA	CAG	GAA	CAA	ACA	AGA	CTT	AAT
	GGA	C GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u>	SEQ II	NO:	15)
YLR440c-S1	TTG	CCA	AGA	AAA	TTG	CAG	TAA	AAA	TGT	TGG	AAG	AGC
	AAC	T CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ I	D NO:	<u>16)</u>	
YLR440c-S2	GCT	CCA	ATT	CTA	GTG	TGC	TCC	ATT	GCG	ATG	TAA	CAA
	TTT	C GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u>	SEQ II	NO:	<u>17)</u>

Block

Gene deletions	on	chro	noso	me 6								
Name	Sequ	ı nce	5'-	-3'		_						
YFL024c-S1	TGA	TGA	ATT	TTT	CTG	GGT	TAT	AGA	AGA	GTT	CTG	TTT
	CGC	T CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	NO:	18)	
YFL024c-S2	ACA	CCT	TCA	AAC	GCT	ATA	GAG	ATC	AAT	GAC	GGT	TCG
	CAT	A GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (S	SEQ II	NO:	20)
YFR003c-S1	TGT	GGA	AGA	GGT	TCC	CGC	AGT	TTT	GCA	GCT	TCG	AGC
	AAC	TCA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	ONO:	21)	
YFR003c-S2	ATC	TTC	TTT	GTC	TAC	GTT	CGT	TAA	AGT	CAA	GAT	CCT
	TCT	C GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u> §	SEQ II	NO:	22)
YFR027w-S1	AAT	GAA	AGC	TAG	GAA	ATC	GCA	GAG	AAA	AGC	GGG	CAG
	TAA	A CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ II	NO:	23)	
YFR027w-S2	AAT	TTG	GTT	GCG	ATA	CCC	AAC	TTC	CTT	GCT	GTC	CTG
	CAC	A GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (S	SEQ II	NO:	24)
YFR042w-S1	AGT	TTG	CAC	CAA	TGG	CAA	TAT	GCC	TGT	GAT	AAA	GAT
	AAG	G CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ II	ONO:	<u>25)</u>	
YFR042w-S2	CAT	GGA	AGT	TAT	TTG	GTT	GCT	TAG	ATT	CCA	CGG	GTT
	CAA	A GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u> §	SEQ II	NO:	26)

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Gene deletions	on	chro	moso	me 9								
Name	S qu	ence	∍ 5'-	-3'								
YIL109c-S1	TGT	CTC	ATC	ACA	AGA	AAC	GTG	TTT	ACC	CAC	AAG	CTC
	AGC	T CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ II	ONO:	27)	
YIL109c-S2	TCA	TGA	TTT	GTA	AGA	ATT	CTC	TGT	AAC	TTT	CGT	TAT
	TCA	A GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u>	SEQ II	NO:	29)
YIL091c-S1	AGT	GAC	AGT	TCT	GTG	AGG	GAA	AAG	AAT	GAT	AAT	TTC
	CGT	G CA	GCT	GAA	GCT	TCG	TAC	GC (SEQ II	ONO:	30)	
YIL091c-S2	CAT	TGT	AAA	ATT	CAG	GAT	TGT	TTG	GAG	GCT	TAT	AAA
	AAA	C GC	ATA	GGC	CAC				TG (
YIL083c-S1	ACC	TCT	ACC	CGT	GCT	CAA	CAG	ACC	TCA	AAT	TCA	TAC
	GTC	TCA			GCT		TAC		SEQ I			
YIL083c-S2	CGA	TGA			GGA				CTT			
	GGT						TGG			SEQ II		
YIL019w-S1	TTC				CTT				AGT			CTT
	t				GCT				SEQ II			
YIL019w-S2	TAG		CGG	GAT					TTA			
	СТС			GGC					TG <u>(</u>			
YIL104c-S1	TCC	CCT	TAC	TAT					TTT			GAA
	 								SEQ II			
YIL104c-S2	CCT	GAT	ACC	TGT					CTT			
	GCT								TG (
YIR010w-S1	ATG		CTG	_					GTC			ACG
WTD010 G0	 	CCA					TAC TTG		SEQ II	ACT		TCT
YIR010w-S2	CAA		GCT	CAT		CTT						
VTD015 C1	GCG	CGC	ATA CAG		CAC	TTC	TGG		TTA	SEQ II		
YIR015w-S1	TAC		GCT		GCT	TCG			SEO II		41)	CIC
YIR015w-S2	ATT		AGT	TTT				<u>~</u>	AAC			ጥርር
IIIOIJWDZ	CAT								TG (S			

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Gene deletions	on	chron	nosor	n 16	5							
Name	Sequ	ience	5'-	-3'								
YPL233w-S1	ATG	TCA	CAA	GGT	CAG	TCC	AAA	AAA	CTG	GAC	GTA	ACT
	GTT	G CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	NO:	43)	
YPL233w-S2	CAA	TCC	TCC						AAG	CGC	TTG	ACC
	TTT	TGC	ATA		CAC		TGG			SEQ II		
YPL146c-S1	TCC	AAC	TAA				GAA		ATC	TCA		CAA
		G CA				TCG		<u> </u>	SEQ II			
YPL146c-S2	TTT		GTC	CTT	ATG				TTC			TTT
	CTG		ATA		CAC		TGG			SEQ II		
YPL126w-S1	ATG	ACG				GGT	_	-	CAG			CTG
VDI 106 00	TCA	G CA		GAA ACT			TAC		SEQ II TCA			GTA
YPL126w-S2	TAT	GTT	AAT		_	ATC						
YPL093w-S1	CAA	AGC GAT	TAC		CAC		TGG AGC	-		SEQ II TAT	GCG	
IPLU93W-BI	AGT	TCA		GAA			TAC		SEQ II			IAA
YPL093w-S2	CGG	AAA	TCT	GTC	TTA				CGC	•		СТС
1FD095W-52	AAT		ATA				TGG		TG (S			
YPL063w-S1	TTC	AAG	CAT				TTT		CAA			
112003# 51	GTT		GCT		_	TCG			SEQ II			
YPL063w-S2	GGA	TTC		AAT	CTT				СТТ			TTC
	AAA			GGC		_	TGG		TG (5	SEQ II	NO:	53)
YPL024w-S1	ATG	TCT	TTT	TCA		ATC			CAG			
	GAT	G CA	GCT	GAA	GCT		TAC	GC (SEQ II	ONO:	54)	
YPL024w-S2	ACT	TGT	GAG	TCC	TTC	AAT	ATG	AAA	ACG	CCC	CTA	TTG
	AAC	A GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (8	SEQ II	NO:	<u>55)</u>
YPL020c-S1	TCA	GTT	GAA	GTA	GAT	AAG	CAC	CGG	AAC	ACA	CTA	CAG
	TAT	C CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	ONO:	56)	
YPL020c-S2	TCG	GTT	AAA	ATC	AAA	TGG	GCA	ATA	AAT	CTT	CTC	ATC
	CTA	A GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (8	SEQ II	NO:	<u>57)</u>
YPL012w-S1	ATG	GAT	CAA	GAC	AAA	GTT	GCT	TTT	CTT	TTA	GAG	CTG
	GAG	G CA	GCT			TCG	TAC	GC <u>(</u>	SEQ II	ONO:	58)	
YPL012w-S2	ATT		ACT		GAC	CTT	TCT	TAT		GTT		CAA
									TG (S			
YPL007c-S1									TCT			GCG
									SEQ II			
YPL007c-S2									ATT			
WDD 0 4 0									TG (S			
YPR048w-S1									ATC			GGA
VDD 0.4 0 C2	·								SEQ II			a Cm
YPR048w-S2	AAT								TG (S			
YPR072w-S1	+								ATA			
IFRU/ZW-SI				_					SEQ II			AGC
VDD 0.7.250- 6.2	T								CTT			ጥጥር
YPR072w-S2	AGA TTT								TG (
	TIT	AGC	AIA	<u> </u>	CAC	IAG	100	AIC	10 (3	عدر الم	140:	30)

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Name	Seq	uence	3 5'-	-3'								
YPR082c-S1	CTT	CGA	TTG	CTG	AAA	GAG	TAA	GGA	ACT	TTG	CAG	TT
	TTT	A CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ I	D NO:	<u>67)</u>	
YPR082c-S2	CAA	TAA	AGT	TCA	ACT	TGT	TGT	TGI	TCC	CTG	TAC	CAF
	AAT	C GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (SEQ II	NO:	68)
YPR085c-S1	CTG	TAC	ATT	CTT	TCG	AAA	GAC	TCC	ATG	CTG	CGA	ΑTΊ
	TTT	G CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ I	D NO:	69)	
YPR085c-S2	TCC	CAC	TTT	ATA	GTT	ATG	GGA	ттт	CGA	GCT	GGA	TTC
	GGT	A GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u>	SEQ II	NO:	70)
YPR105c-S1	AGC	TCG	ATC	ATC	GAG	GGC	CAA	TTG	TCT	AAA	AAT	CTA
	GCA	A CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ I	D NO:	72)	
YPR105c-S2	CTG	TGT	TCT	ATC	AAT	CTT	CAT	ATI	TCT	AGC	TTT	LAA
	TCT	TGC	ATA	GGC	CAC	TAG	TGG	ATC	TG (SEQ II	NO:	73)
YPR112c-S1	CAT	TGT	CAA	GGG	TTT	GCC	CGT	CTA	TCT	AAC	AGA	TGA
	TAA	T CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ I	D NO:	74)	
YPR112c-S2	GAA	ACC	TTC	GTT	TTC	TTC	ATC	ATC	CAC	ATC	CAG	TTT
	CTT	TGC	ATA	GGC	CAC	TAG	TGG	ATC	TG (SEQ II	NO:	76)
YPR137w-S1	ATG	TCA	GAT	GTT	ACC	CAA	CAG	AAA	AAG	AGG	AAA	AGA
	TCC	A CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ I	D NO:	<u>75)</u>	
YPR137w-S2	AAA	AGC	CTG	TTT	GGT	CAA	TGA	CAG	CTG	AAT	ATA	TAC
	CAT	TGC	AT <i>P</i>	A GG	C CA	C TA	G T	GG A	TC I	'G <u>(s</u> 1	EQ ID	NO:
	180)											
YPR143w-S1	ATG	GGC	TCC	AAG	CAC	AGA	GTA	GAC	ACT	AAG	GAT	AAG
	AAA	A CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ I	D NO:	77)	
YPR143w-S2	TTC	ATT	GTC	GCT	TCC	TGC	GGC	AGC	TTT	AAC	TAA	ATC
	CAA	A GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (SEQ II	NO:	78)
YPR144c-S1	TTC	CAG	AAA	ATG	TTA	CTC	AAT	TGG	AAG	AAG	ATG	AGA
	CAG	A CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ I	D NO:	79)	
YPR144c-S2	CCA	TGC	TAC	CCC	AGG	CAA	GTA	GAC	GTT	ACC	TTG	GGA
	TGA	C GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u>	SEQ II	NO:	80)
YPR169w-S1	TTT	TAC	ATC	CTG	AAC	TGC	CCA	TTA	TAA	TAA	CTG	GCI
	TTG	G CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ I	D NO:	82)	
YPR169w-S2	CTT	CTT	GAT	CCC	ATG	CTC	ATA	CAG	GTC	CTT	TTT	TTT
	GTT	G GC	АТА	GGC	CAC	TAG	TGG	ATC	TG (SEQ II	NO:	83)

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